



EXHIBIT A

SEQ5	1	MAFWTQLMLLLWKNFMYRRQPVQLLVELLWPLFLFFILVAVRHSHPPLE	50
SEQ6	1	PPLE	4

SEQ5	51	HHECHFPNKPPLPSAGTPWLIQGLICNVNNNTCFPQLTPGEEPGRLSNFNDS	100
SEQ6	5	HHECHFPNKPPLPSAGTPWLIQGLICNVNNNTCFPQLTPGEEPGRLSNFNDS	54

SEQ5	101	LVSRLADARTVLGGASAHTLAGLGKLIATLRAARSTAQPQPTKQSPLE	150
SEQ6	55	LVSRLADARTVLGGASAHTLAGLGKLIATLRAARSTAQPQPTKQSPLE	104

SEQ5	151	PPMLDVAELLTSLLRTESLGLALGQAQEPLHSLLEAAEDLAQELLALRSL	200
SEQ6	105	PPMLDVAELLTSLLRTESLGLALGQAQEPLHSLLEAAEDLAQELLALRSL	154

SEQ5	201	VELRALLQRPRTSGPPELLSEALCSVRGPSSTVGPSLNWYEASDLMELV	250
SEQ6	155	VELRALLQRPRTSGPPELLSEALCSVRGPSSTVGPSLNWYEASDLMELV	204

SEQ5	251	GQEPEPALPDSSLSPACSELIGALDSHPLSRLWRRLKPLILGKLLFAPD	300
SEQ6	205	GQEPEPALPDSSLSPACSELIGALDSHPLSRLWRRLKPLILGKLLFAPD	254

SEQ5	301	TPFTRKLMAQVNRTFEELTLLRDVREVWEMLGPRIFTFMNDSSNVAMLQR	350
SEQ6	255	TPFTRKLMAQVNRTFEELTLLRDVREVWEMLGPRIFTFMNDSSNVAMLQR	304

SEQ5	351	LLQMDEGRRQPRPGGRDHMEALRSFLDPGSGGGYSWQDAHADVGHLVGTL	400
SEQ6	305	LLQMDEGRRQPRPGGRDHMEALRSFLDPGSGGGYSWQDAHADVGHLVGTL	354

SEQ5	401	GRVTECLSLDKLEAAPSEAALVSRALQLLAEHRFWAGVVFLGPEDSSDPT	450
SEQ6	355	GRVTECLSLDKLEAAPSEAALVSRALQLLAEHRFWAGVVFLGPEDSSDPT	404

SEQ5	451	EHPTPDLGPGHVRIKIRMDIDVVTRTNKIRDRFWDPGPAADPLTDLRYVW	500
SEQ6	405	EHPTPDLGPGHVRIKIRMDIDVVTRTNKIRDRFWDPGPAADPLTDLRYVW	454

SEQ5	501	GGFVYLQDLVERAARVLVLSGANPRAGLYLQQMPYPCYVDDVFLRVLSRSL	550
SEQ6	455	GGFVYLQDLVERAARVLVLSGANPRAGLYLQQMPYPCYVDDVFLRVLSRSL	504

SEQ5	551	PLFLTLAWIYSVTLTVKAVVREKETRLRDTMRAMGLSRAVLWLGWFLSCL	600
SEQ6	505	PLFLTLAWIYSVTLTVKAVVREKETRLRDTMRAMGLSRAVLWLGWFLSCL	554

SEQ5	601	GPFLLSAALLVLVLKLGDLILPYSHPGVVFLAFAVATVTQSFLLSAFF	650
SEQ6	555	GPFLLSAALLVLVLKLGDLILPYSHPGVVFLAFAVATVTQSFLLSAFF	604

SEQ5	651	SRANLAAACGGLAYFSLYLPYVLCVAWRDRLPAGGRVAASLLSPVAFGFG	700
SEQ6	605	SRANLAAACGGLAYFSLYLPYVLCVAWRDRLPAGGRVAASLLSPVAFGFG	654

SEQ5	701	CESLALLEEQGEGAQWHNVGTRPTADVFSLAQVSGLLLDAALYGLATWY	750
SEQ6	655	CESLALLEEQGEGAQWHNVGTRPTADVFSLAQVSGLLLDAALYGLATWY	704

SEQ5	751	LEAVCPGQYGIPEPWNPFPFRSYWCGPRPPKSPAPCPTPLDPKVLVEEAP	800
SEQ6	705	LEAVCPGQYGIPEPWNPFPFRSYWCGPRPPKSPAPCPTPLDPKVLVEEAP	754

SEQ5	801	PGLSPGVSVRSLEKRFPGSPQPALRGLSLDFYQGHITAFLGHNGAGKTTT	850
SEQ6	755	PGLSPGVSVRSLEKRFPGSPQPALRGLSLDFYQGHITAFLGHNGAGKTTT	804

SEQ5	851	LSILSGLFPPSGGSAFILGHDRVSSMAAIRPHLGVCPOQYNVLFDMLTVDE	900
SEQ6	805	LSILSGLFPPSGGSAFILGHDRVSSMAAIRPHLGVCPOQYNVLFDMLTVDE	854

SEQ5	901	HVWFYGRLKGLSAAVGPEQDRLLQDVGLVKQSVQTRHLSGGMQRKLSV	950
SEQ6	855	HVWFYGRLKGLSAAVGPEQDRLLQDVGLVKQSVQTRHLSGGMQRKLSV	904

SEQ5	951	AIAFVGGSQVVIILDEPTAGVDPASRRGIWELLKYREGRTLILSTHHDE	1000
SEQ6	905	AIAFVGGSQVVIILDEPTAGVDPASRRGIWELLKYREGRTLILSTHHDE	954

SEQ5	1001	AELLGDRVAVVAGGRLCCGSPLFLRRHLGSGYYLTLVKARLPLTTNEKA	1050
SEQ6	955	AELLGDRVAVVAGGRLCCGSPLFLRRHLGSGYYLTLVKARLPLTTNEKA	1004

SEQ5	1051	DTDMEGSVDTRQEKKNGSQGSRVGTPQLLALVQHWVPGARLVEELPHELV	1100
SEQ6	1005	DTDMEGSVDTRQEKKNGSQGSRVGTPQLLALVQHWVPGARLVEELPHELV	1054

SEQ5	1101	LVLPTYGAHDGSFATLFRELDTRLAELRLTGYYGDISDTSLEEIFLKVVVEEC	1150
SEQ6	1055	LVLPTYGAHDGSFATLFRELDTRLAELRLTGYYGDISDTSLEEIFLKVVVEEC	1104

SEQ5	1151	AADTDMEDGSCGQHLCTGIAGLDVTLRLKMPHQETALENGEPAGSAPETD	1200
SEQ6	1105	AADTDMEDGSCGQHLCTGIAGLDVTLRLKMPHQETALENGEPAGSAPETD	1154

SEQ5	1201	QGSGPDAVGRVQGWALTRQQLQALLLKRFLLARRSRRGLFAQIVLPALFV	1250
SEQ6	1155	QGSGPDAVGRVQGWALTRQQLQALLLKRFLLARRSRRGLFAQIVLPALFV	1204

SEQ5	1251	GLALVFSLIVPPFGHYPALRLSPTMYGAQVSSFFSEDAPGDPGRARLLEAL	1300
SEQ6	1205	GLALVFSLIVPPFGHYPALRLSPTMYGAQVSSFFSEDAPGDPGRARLLEAL	1254

SEQ5	1301	LQEAGLEEPPVQHSSHRFSAPEVPAEVAKVLASGNWTPESPSPACQCSQP	1350
SEQ6	1255	LQEAGLEEPPVQHSSHRFSAPEVPAEVAKVLASGNWTPESPSPACQCSQP	1304

SEQ5 1351 GARRLLPDCPAAAGGPPPPQAVTGSGEVVQNLTGRNLSDFLVKTYPRLVR 1400
 SEQ6 1305 GARRLLPDCPAAAGGPPPPQAVTGSGEVVQNLTGRNLSDFLVKTYPRLVR 1354

 SEQ5 1401 QGLKTKKWWVNEVRYGGFSLGGRDPGLPSQELGRSVEELWALLSPLPGGA 1450
 SEQ6 1355 QGLKTKKWWVNEVRYGGFSLGGRDPGLPSQELGRSVEELWALLSPLPGGA 1404

 SEQ5 1451 LDRV LKNLTAWAHSLDAQDSLKIWFNNKGWHSMVAFVN RASNAILRAHLP 1500
 SEQ6 1405 LDRV LKNLTAWAHSLDAQDSLKIWFNNKGWHSMVAFVN RASNAILRAHLP 1454

 SEQ5 1501 PGPARH AHSITTLN HPLN LTKEQLSEAALMASSVDL VSICV VFAM SFVP 1550
 SEQ6 1455 PGPARH AHSITTLN HPLN LTKEQLSEAALMASSVDL VSICV VFAM SFVP 1504

 SEQ5 1551 ASFTL VL LIEERV TRAKH LQLM GGLSPTLYWL GNFLW DMCN YL VPACIV VL 1600
 SEQ6 1505 ASFTL VL LIEERV TRAKH LQLM GGLSPTLYWL GNFLW DMCN YL VPACIV VL 1554

 SEQ5 1601 IFLAFQQRAYVAPANLPALLLLLLY GWSITPLM PASF FFSVP STAY VV 1650
 SEQ6 1555 IFLAFQQRAYVAPANLPALLLLLLY GWSITPLM PASF FFSVP STAY VV 1604

 SEQ5 1651 LTCINL FIGINGSMATF VLELFSDQKLQEV S RILKQVFLIFPHFC LGRGL 1700
 SEQ6 1605 LTCINL FIGINGSMATF VLELFSDQKLQEV S RILKQVFLIFPHFC LGRGL 1654

 SEQ5 1701 IDMVRN QAMADA FERL GDRQ FQSPLRWEVVGKNLLAMVIQGPLFLLFTLL 1750
 SEQ6 1655 IDMVRN QAMADA FERL GDRQ FQSPLRWEVVGKNLLAMVIQGPLFLLFTLL 1704

 SEQ5 1751 LQHRSQ LLPQPR VRS LPLLGE EDEDVAR E RER VVQG ATQGDV LVL RNLTK 1800
 SEQ6 1705 LQHRSQ LLPQPR VRS LPLLGE EDEDVAR E RER VVQG ATQGDV LVL RNLTK 1754

 SEQ5 1801 VYRGQRMPA VDRLCLGIPPGECFG LGVNGAGKTSTFRMVTGDT LASRGE 1850
 SEQ6 1755 VYRGQRMPA VDRLCLGIPPGECFG LGVNGAGKTSTFRMVTGDT LASRGE 1804

 SEQ5 1851 AVL A GH SVA REPSA A HLSM GYCP QSDA I FELL TGREH LELLAR LRGV PEA 1900
 SEQ6 1805 AVL A GH SVA REPSA A HLSM GYCP QSDA I FELL TGREH LELLAR LRGV PEA 1854

 SEQ5 1901 QVAQ TAGS GLAR LGL SWYAD RPAG TYSG GNKR KLATA LAL VGDPA VFLD 1950
 SEQ6 1855 QVAQ TAGS GLAR LGL SWYAD RPAG TYSG GNKR KLATA LAL VGDPA VFLD 1904

 SEQ5 1951 EPTTGMDPSARRFLWNSLLAVVREGRS VMLTSHSMEECEALCSRLAIMVN 2000
 SEQ6 1905 EPTTGMDPSARRFLWNSLLAVVREGRS VMLTSHSMEECEALCSRLAIMVN 1954

 SEQ5 2001 GRFRCLGSPQHLKGRFAAGHTLRLRVPAARSQPA AAFVAAEFPGSELREA 2050
 SEQ6 1955 GRFRCLGSPQHLKGRFAAGHTLRLRVPAARSQPA AAFVAAEFPGSELREA 2004

SEQ5 2051 HGGRLRFQLPPGGRCALARVFGELAVHGAEHGVEDFSVSQTMLEEVFLYF 2100
SEQ6 2005 HGGRLRFQLPPGGRCALARVFGELAVHGAEHGVEDFSVSQTMLEEVFLYF 2054

SEQ5 2101 SKDQGKDEDTEEQKEAGVGVDPAAPGLQHPKRVSQFLDDPSTAETVL 2146
SEQ6 2055 SKDQGKDEDTEEQKEAGVGVDPAAPGLQHPKRVSQFLDDPSTAETVL 2100

EXHIBIT B

SEQ6	1	PPLEHHECHFPNKPLPSAGTVPWLQGLICNVNNNTCFPQLTPGEEPGRLSN	50
SEQ8	1		0
SEQ6	51	FNDSLVSRLADARTVLGGASAHRRTLALGKLIATLRAARSTAQPQPTKQ	100
SEQ8	1	MVCLGTGQSAGPLVSVQNHCPPCGLS-	26
		* * . * . . . *	
SEQ6	101	SPLEPPMLDVAELLTSLLRTESLGLALGQAQEPLHSLLEAAEDLAQELLA	150
SEQ8	27	-PQ-----ESLGLALGQAQEPLHSLLEAAAGDLAQELLA	58
	*	*****	*****
SEQ6	151	LRSI VELRALLQ QPRGTSGPLELLSEALCSVRGPSSTVGPSLNWYEASDL	200
SEQ8	59	LRSI VELRALLQ QPRGTSGPLELLSEALCSVRGPSSTVGPSLNWYEASDL	108
		*****	*****
SEQ6	201	MELVGQEPE SALPDSSL SPACSELIGALD SH PLSRL W RRLKPLILGKLL	250
SEQ8	109	MELVGQEPE SALPDSSL SPACSELIGALD SH PLSRL W RRLKPLILGKLL	158
		*****	*****
SEQ6	251	FAPDTPFTRKLMAQVNRTFEELTLLRDVREVWEMLGPRIFTFMNDSSNVA	300
SEQ8	159	FAPDTPFTRKLMAQVNRTFEELTLLRDVREVWEMLGPRIFTFMNDSSNVA	208
		*****	*****
SEQ6	301	MLQRLLQM QDEGRRQ PRPG RDH MEALRSFL D PGSGGYSWQDAHADV GHL	350
SEQ8	209	MLQRLLQM QDEGRRQ PRPG RDH MEALRSFL D PGSGGYSWQDAHADV GHL	258
		*****	*****
SEQ6	351	VGTLGRVTECLSLDKLEAAPSEAALVS R ALQ L A H RFWAGVVFLGPEDS	400
SEQ8	259	VGTLGRVTECLSLDKLEAAPSEAALVS R ALQ L A H RFWAGVVFLGPEDS	308
		*****	*****
SEQ6	401	SDPTEHPTPD L PGHVR I KIRMDIDVV T RTN K IRDRF W D P GAADPL T DL	450
SEQ8	309	SDPTEHPTPD L PGHVR I KIRMDIDVV T RTN K IRDRF W D P GAADPL T DL	358
		*****	*****
SEQ6	451	RYVWGGFVYLQDLVERA AV RV L SGANP R AGLYLQQMPYPCYVDDVFLRVL	500
SEQ8	359	RYVWGGFVYLQDLVERA AV RV L SGANP R AGLYLQQMPYPCYVDDVFLRVL	408
		*****	*****
SEQ6	501	SRS L PL F TLAWIYSVT L TVKAVVRE K E T RL R DTM R AMGLSRAV L WL G WF	550
SEQ8	409	SRS L PL F TLAWIYSVT L TVKAVVRE K E T RL R DTM R AMGLSRAV L WL G WF	458
		*****	*****
SEQ6	551	LSCLGP F LLSA A LLV L V L K L G D I L P Y SHPGVVFL F LA A FA V AT V TQS F LL	600
SEQ8	459	LSCLGP F LLSA A LLV L V L K L G D I L P Y SHPGVVFL F LA A FA V AT V TQS F LL	508
		*****	*****
SEQ6	601	SAFFSRANLAAACGG L AYFS L PYV L CV A WR D RLP A GG R V A AS L SP V A	650
SEQ8	509	SAFFSRANLAAACGG L AYFS L PYV L CV A WR D RLP A GG R V A AS L SP V A	558
		*****	*****

SEQ6	651	FGFGCESLALLEEQGEGAQWHNVGTRPTADVFSLAQVSGLLLDAALYGL	700
SEQ8	559	FGFGCESLALLEEQGEGAQWHNVGTRPTADVFSLAQVSGLLLDAALYGL	608

SEQ6	701	ATWYLEAVCPGQYGIPEPWNPFRRSYWCGRPPKSPAPCPTPLDPKV	750
SEQ8	609	ATWYLEAVCPGQYGIPEPWNPFRRSYWCGRPPKSPAPCPTPLDPKV	658

SEQ6	751	EEAPPGLSPGVSVRSLEKRFPGSPQPALRGLSLDFYQGHITAFLGHNGAG	800
SEQ8	659	EEAPPGLSPGVSVRSLEKRFPGSPQPALRGLSLDFYQGHITAFLGHNGAG	708

SEQ6	801	KTTLISLGLFPPSGGSAFILGHDRSSMAAIRPHLGVCQYNVLFDML	850
SEQ8	709	KTTLISLGLFPPSGGSAFILGHDRSSMAAIRPHLGVCQYNVLFDML	758

SEQ6	851	TVDEHWFYGRLKGLSAAVVGPEQDRLLQDVGLVSKQSVQTRHLSGGMQR	900
SEQ8	759	TVDEHWFYGRLKGLSAAVVGPEQDRLLQDVGLVSKQSVQTRHLSGGMQR	808

SEQ6	901	KLSVAIAFVGGSQVVILDEPTAGVDPASRRGIWELLKYREGRTLILSTH	950
SEQ8	809	KLSVAIAFVGGSQVVILDEPTAGVDPASRRGIWELLKYREGRTLILSTH	858

SEQ6	951	HLDEAELLGDRVAVVAGGRLCCCGSPLFLRRHLGSGYYLTIVKARLPLTT	1000
SEQ8	859	HLDEAELLGDRVAVVAGGRLCCCGSPLFLRRHLGSGYYLTIVKARLPLTT	908

SEQ6	1001	NEKADTDMEGSVDRQEKKNQSGSRVGTQPLLALVQHWVPGARLVEELP	1050
SEQ8	909	NEKADTDMEGSVDRQEKKNQSGSRVGTQPLLALVQHWVPGARLVEELP	958

SEQ6	1051	HELVLVLPYTGAHDGSFATLFRELDTRLAELRLTGYGISDTSLEEIFLV	1100
SEQ8	959	HELVLVLPYTGAHDGSFATLFRELDTRLAELRLTGYGISDTSLEEIFLV	1008

SEQ6	1101	VEECAADTDMEDGSCGQHLCTGIAGLDVTLRLKMPQETALENGEPAGSA	1150
SEQ8	1009	VEECAADTDMEDGSCGQHLCTGIAGLDVTLRLKMPQETALENGEPAGSA	1058

SEQ6	1151	PETDQGSGPDAVGRVQGWALTRQQLQALLKRFLLARRSRRGLFAQIVLP	1200
SEQ8	1059	PETDQGSGPDAVGRVQGWALTRQQLQALLKRFLLARRSRRGLFAQIVLP	1108

SEQ6	1201	ALFVGLALVFSLIVPPFGHYPALRLSPTMYGAQVSFFSEDAPGDPGRARL	1250
SEQ8	1109	ALFVGLALVFSLIVPPFGHYPALRLSPTMYGAQVSFFSEDAPGDPGRARL	1158

SEQ6	1251	LEALLQEAGLEEPPVQHSSHRFSAPEVPAEVAKVLASGNWTPESPSPACQ	1300
SEQ8	1159	LEALLQEAGLEEPPVQHSSHRFSAPEVPAEVAKVLASGNWTPESPSPACQ	1208

SEQ6	1301	CSQPGARRLLPDCPAAAGGPPPPQAVTGSGEVVQNLGRNLSDFLVKTYP	1350
SEQ8	1209	CSQPGARRLLPDCPAAAGGPPPPQAVTGSGEVVQNLGRNLSDFLVKTYP	1258

SEQ6 1351 RLVRQGLTKKWWNEVRYGGFSLGGRDPGLPSGQELGRSVEELWALLSPL 1400
 SEQ8 1259 RLVRQGLTKKWWNEVRYGGFSLGGRDPGLPSGQELGRSVEELWALLSPL 1308

 SEQ6 1401 PGGALDRVVLKNLTAWAHSLDAQDSLKIWFNNKGWHSMVAFVNRAASNAILR 1450
 SEQ8 1309 PGGALDRVVLKNLTAWAHSLDAQDSLKIWFNNKGWHSMVAFVNRAASNAILR 1358

 SEQ6 1451 AHLPPGPARHAHSITTLNHPNLTKEQLSEAALMASSVDVLVSICVVFAM 1500
 SEQ8 1359 AHLPPGPARHAHSITTLNHPNLTKEQLSEAALMASSVDVLVSICVVFAM 1408

 SEQ6 1501 SFVPASFTLVVIEERVTRAKHLQLMGLSPTLYWLGNFLWDMCNLYLPAC 1550
 SEQ8 1409 SFVPASFTLVVIEERVTRAKHLQLMGLSPTLYWLGNFLWDMCNLYLPAC 1458

 SEQ6 1551 IVVLIIFLAFQQRAYVAPANLPALLLLLLYGWSITPLMPASFFFSVPST 1600
 SEQ8 1459 IVVLIIFLAFQQRAYVAPANLPALLLLLLYGRQAVPVTALGGGRQEP-- 1506
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 SEQ6 1601 AYVVLTCINLFIGINGSMATFVLELFSDQKLQEVSRILKQVFLIFPHFCL 1650
 SEQ8 1507 ----- 1506

 SEQ6 1651 GRGLIDMVRNQAMADAFLRQFQSPRLWEVVGKNLLAMVIQGPLFLL 1700
 SEQ8 1507 ----- 1506

 SEQ6 1701 FTLLLQHRSQLLPQPRVRSLPLLGEEDEDVARERERVVQGATQGDVLVLR 1750
 SEQ8 1507 ---LGHGDTGAPLP----- 1517
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 SEQ6 1751 NLTKVYRGQRMPAVDRLCLGIPPGEFCGLLGVNGAGKTSTFRMVTGDTLA 1800
 SEQ8 1518 ----- 1517

 SEQ6 1801 SRGEAVLAGHSVAREPSAAHLSMGYCPQSDAIFELLTGREHLELLARLRG 1850
 SEQ8 1518 ----- 1517

 SEQ6 1851 VPEAQVAQTAGSGLARLGLSWYADRPAGTYSGGNKRKLATALALVGDPAV 1900
 SEQ8 1518 ----- 1517

 SEQ6 1901 VFLDEPTTGMDPSARRFLWNSLLAVVREGRSVMLTSHSSEECEALCSRLA 1950
 SEQ8 1518 ----- 1517

 SEQ6 1951 IMVNNGRFRCGLSPQHLKGRFAAGHTLTLRVPAARSQPAAFVAAEFPGSE 2000
 SEQ8 1518 -----SLHTTAAAPKPTPATAQGEVSATP 1541
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 SEQ6 2001 LREAHGGRLRFQLPPGGRCALARVFGELAVHGAEHGVEDFSVSQTMLEEV 2050
 SEQ8 1542 GRGGRGCGSP 1550
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 SEQ6 2051 FLYFSKDQGKDEDTEEQKEAGVGVDPAPGLQHPKRVSQFLDDPSTAETVL 2100



EXHIBIT C

SEQ2	1 MAFCTQLMLLLWKNYTYRRRQPIQLLVELLWPLFLFFFILVAVRHSHPPLE	50
SEQ5	1 MAFWTQLMLLLWKNFMYRRRQPVQLLVELLWPLFLFFFILVAVRHSHPPLE	50
	***** . *****.*****.*****.*****.*****.*****.*****.*****	
SEQ2	51 HHECHFPNKPPLPSAGTVPWLQGLVCNVNNSCFQHPTPGEKPGVLSNFKDS	100
SEQ5	51 HHECHFPNKPPLPSAGTVPWLQGLICNVNNNTCPQLTPGEEPGRLSNFNDS	100
	*****.*****.*****.*****.*****.*****.*****.*****.*****	
SEQ2	101 LISRLLADTRTVLGGHSIQDMLDALGKLIPVLRAGGGARPQESDQPTSQ	150
SEQ5	101 LVSRLADARTVLGASAHLRTLAGLGKLIATLRAARSTAQPQPTKQSPLE	150
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SEQ2	151 G---SVTKLLEKILQRASLDPVLGQAQDSMRKFSDAIRDLAQELLTLPSL	197
SEQ5	151 PPMLDVAELLTSLRTESLGLALGQAQEPLHSLLEAAEDLAQELLALRSL	200
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SEQ2	198 MELRALLRRPRGSAGSLELVSEALCSTKGPPSPGGLSLNWYEANQLNEFM	247
SEQ5	201 VELRALLQRPRGTSGPELLSEALCSVRGPSSTVGPSLNWYEASDLMELV	250
	.*****.*****..* ***.***** .***** * ***** * ***** * .	
SEQ2	248 GPEVAPALPDNSLSPACSEFVGTLDDHPVSRLLWRRLKPLILGKILFAPD	297
SEQ5	251 GQEPEALPDSSLSPACSELIGALDSHPLSRLLWRRLKPLILGKLLFAPD	300
	* * *** ***** .*.** * .*****.*****.*****.*****	
SEQ2	298 TNFTRKLMAQVNQTFEELALLRDHELWGVLPQIFNFMDSTNVAMLQR	347
SEQ5	301 TPFTRKLMQAQNRTFEELLLRDREVWEMLGPRIFTFMNDSSNVAMLQR	350
	* *****.*****.*****.*****. . * . * . * . * . * . * . * .	
SEQ2	348 LLDVGGTGQRQQTPRAQKKLEAIKDFLDPSRGGYSWREAHADMGRLAGIL	397
SEQ5	351 LLQMDEGRRQPRPGGRDHMEALRSFLDPGGGGYSQDAHADVGHVGTL	400
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SEQ2	398 GQMMECVSLDKLEAVPSEEALVSRALELLGERRLWAGIVFLSPEHPLDPS	447
SEQ5	401 GRVTECLSLDKLEAAPSEAALVSRALQLLAEHRFWAGVVFLGPEDSSDPT	450
	*.. * .***** * . * . * . * . * . * . * . * . * . * . * .	
SEQ2	448 ELSSPALSPGHLRFKIRMDIDDVTRTNKIRDKFWDPGAPSADPFMDLRYVW	497
SEQ5	451 EHPTPDLGPGHVRIKIRMDIDVVTRTNKIRDRFWDPGPAADPLTDLRYVW	500
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SEQ2	498 GGFVYLQDLLEQAAVRVLGGGNSRTGLYLOQQMPHPCYVDDVFLRVLSRSL	547
SEQ5	501 GGFVYLQDLVERAAVRVLSGANPAGLYLOQQMPYPCYVDDVFLRVLSRSL	550
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SEQ2	548 PLFLTLAWIYSVALTVKAVVREKETRLRETMRAMGLSRAVLWLGWFLSCL	597
SEQ5	551 PLFLTLAWIYSVTLTVKAVVREKETRLRDTMRAMGLSRAVLWLGWFLSCL	600
	*****.*****.*****.*****.*****.*****.*****.*****.*****	
SEQ2	598 GPFLVSAALLVLVLKLGNIIPYSHPVVIFLFLAAFAVATVAQSFLSAFF	647
SEQ5	601 GPFLLSAALLVLVLKLGDIIPYSHPGVVFLFLAAFAVATVTQSFLSAFF	650
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SEQ2	648	SRANLAAACGGLAYFALYLPYVLCVAWRERLHLGGLLAASLLSPVAFGFG	697
SEQ5	651	SRANLAAACGGLAYFSLYLPYVLCVAWRDRLPAGGRVAASLLSPVAFGFG	700

SEQ2	698	CESLALLEEQGDGAQWHNLGTGPAEDVFSLAQSAFLLLDAVIYGLALWY	747
SEQ5	701	CESLALLEEQGEGAQWHNVGTRPTADVFSLAQVSGLLLDAALYGLATWY	750

SEQ2	748	LEAVCPGQYGIPEPWNPFPRRSYWCGPGPPKSSVLAPAPQDPKVLVEPP	797
SEQ5	751	LEAVCPGQYGIPEPWNPFPRRSYWCGPRPPKSPACPPTPLDPKVLVEAP	800

SEQ2	798	LGLVPGVSIRGLKKHFRGCPQPALQGLNLDFYEGHITAFLGHNGAGKTTT	847
SEQ5	801	PGLSPGVSVRSLEKRFPGSPQPALRGLSLDFYQGHITAFLGHNGAGKTTT	850

SEQ2	848	LSILSGLFPPSSGSASI LGHDVQTNMAAIRPHLGICPQYNVLFDMLTVEE	897
SEQ5	851	LSILSGLFPPSSGSAFILGHHDVRSSMAAIRPHLGVC P QYNVLFDMLTVD	900

SEQ2	898	HVWFYGRKGVSAAAMGPERERLIRDVGTLKRDQT RHL SGGMQRKLSV	947
SEQ5	901	HVWFYGRKGLSAAVVGPEQDRLLQDVGLVSKQSVQTRHL SGGMQRKLSV	950

SEQ2	948	AIAFVGGSRVVIMDEPTAGVDPASRRGIWELLKYREGRTLILSTHHLD	997
SEQ5	951	AIAFVGGSQVVI DEPTAGVDPASRRGIWELLKYREGRTLILSTHHLD	1000

SEQ2	998	AELLGDRVAMVAGGS LCCGSPFLRRHLGCGYYLT LVKSSQSLVTHDAK	1047
SEQ5	1001	AELLGDRVAVVAGGR LCCGSPFLRRHLGCGYYLT LVKARLPLTTNEKA	1050

SEQ2	1048	GDSEDPRREKKSDGNGRTSDTAFTRGTSKSNQAPAPGAVPITPSTARIL	1097
SEQ5	1051	DTDMEGSVDTRQE-----KKN-----GSQGSRVGTPQLL	1079

SEQ2	1098	ELVQQHVPAGQLVEDLPHELLLVL PYAGALDGSFAMVFQELDQQLELLGL	1147
SEQ5	1080	ALVQHWVPGARLVEELPHELVLVLPYTGAHDGSFATLFRELDTRLAELRL	1129

SEQ2	1148	TGYGISDTNLEEIFLKVVDAHREGGDSRPQLHLRTCTPQP-----P	1189
SEQ5	1130	TGYGISDTSLEEIFLKVEECAADTD MEDGSCGQHLCTGIAGLDVTLRLK	1179

SEQ2	1190	TGPEASVLENGELAKLVLDPQAPQGLAP-NAAQVQGWTLTCQQLRALLHK	1238
SEQ5	1180	MPPOETALEN GE PAG--SAPETDQGSGPDAVGRVQGWALTRQQLQALLLK	1227

SEQ2	1239	RFLLARRSRRGLFAQVVLPA LFVGLALFFSLIVPPFGQYPPQLSPAMYG	1288
SEQ5	1228	RFLLARRSRRGLFAQIVLPA LFVGLALVFS LIVPPFGHYPALRLSPTMYG	1277

SEQ2	1289	PQVSFFSEDAPGDPNRMKLLEALLGEAGLQEPEPSMQDKDARGSECTHSLAC	1338
SEQ5	1278	AQVSFFSEDAPGDPGRARLLEALLQEAGLEEPPVQHSSHR-----	1317

SEQ2	1339	YFTVPEVPPDVASILASGNWTPESPSPACQCSQPGARRLLPDCPAGAGGP	1388
SEQ5	1318	-FSAPEVPAEVAKVLASGNWTPESPSPACQCSQPGARRLLPDCPAAAGGP	1366
		***** . * . *****	*****
SEQ2	1389	PPPQAVAGLGEVVQNLTGRNVSDLVKTYPQLVRRGLKTKKWWDEVRYGG	1438
SEQ5	1367	PPPQAVTGSGEVVQNLTGRNLSDFLVKTYPRLVRQGLKTKKWWNEVRYGG	1416
		***** . * . *****	*****
SEQ2	1439	FSLGGRDPDLPTGHEVVRTLAEIRALLSPQPGNALDRILNNLTQWALGLD	1488
SEQ5	1417	FSLGGRDPGLPSGQELGRSVEELWALLSPLPGALDRVLKNLTAWAHSLD	1466
		***** . * . *****	*****
SEQ2	1489	ARNSLKIWFNNKGWHAMVAFVN RAN NGLLHALLPSGPVRHAHSITTLNHP	1538
SEQ5	1467	AQDSLKIWFNNKGWHSMVAFVN RASNAILRAHLPNGPARHAHSITTLNHP	1516
		***** . * . *****	*****
SEQ2	1539	LNLTK EQLSEATLIASSV DVLV SICVV FAMS FVPASFTL VLIEERITRAK	1588
SEQ5	1517	LNLTK EQLSEA ALMASSV DVLV SICVV FAMS FVPASFTL VLIEERVTRAK	1566
		***** . * . *****	*****
SEQ2	1589	HLQLVSGLPQ TLW LGNFLW DMCN YL VAVCIVV FIFL AFQQRAYVAPENL	1638
SEQ5	1567	HLQLMGGLSP TLW LGNFLW DMCN YL VPACIVV LIFL AFQQRAYVAPANL	1616
		***** . * . *****	*****
SEQ2	1639	PALL LLLLYGWSITPLMYPASFFF SVPSTAYV VLTCINLFIGINSSMAT	1688
SEQ5	1617	PALL LLLLYGWSITPLMYPASFFF SVPSTAYV VLTCINLFIGINGSMAT	1666
		***** . * . *****	*****
SEQ2	1689	FVLELLSDQNLQEV SRLIKQVFLIFPHFC LGRGLIDMVRNQAMADA FERL	1738
SEQ5	1667	FVLELFSDQKLQEV SRLIKQVFLIFPHFC LGRGLIDMVRNQAMADA FERL	1716
		***** . * . *****	*****
SEQ2	1739	GDKQFQSPLRWDIIGK NLLAMMAQGPLFLLITLLQH RNRLLPQSKP RLL	1788
SEQ5	1717	GDRQFQSPLRWEVVGKNLLAMVIQGPLFLLFTLLQHRSQLLPQPRVRSL	1766
		***** . * . *****	*****
SEQ2	1789	PPLGEEDEDVAQERERVTKGATQGDVLVLRDLTKVYRGQRNP AVDRLCLG	1838
SEQ5	1767	PLLGEEDEDVARERERVQGATQGDVLVLRNLTKVYRGQRMP AVDRLCLG	1816
		***** . * . *****	*****
SEQ2	1839	IPPGECFG LLGVNGAGKTSTFRMVTGDTLPSSGEAVL AGHNV A QERSAAH	1888
SEQ5	1817	IPPGECFG LLGVNGAGKTSTFRMVTGDTLASRGEAVL AGH SVA REPSAAH	1866
		***** . * . *****	*****
SEQ2	1889	RSMGYCPQSDAIFDLLTGREHLELFARL RGVPEA QVA QTALSGL VRLGLP	1938
SEQ5	1867	LSMGYCPQSDAIFELLTGREHLELLARL RGVPEA QVA QTAGS GLARL GLS	1916
		***** . * . *****	*****
SEQ2	1939	SYADRPAGTYSGGNKRKLATALALVGDP AVVFLDEPTTGMDPSARRFLWN	1988
SEQ5	1917	WYADRPAGTYSGGNKRKLATALALVGDP AVVFLDEPTTGMDPSARRFLWN	1966
		***** . * . *****	*****
SEQ2	1989	SLLSVVREGRSVVL TSHS MEECEALCTRLA IVMN GFRCLGSSQHLKGRF	2038
SEQ5	1967	SLLAVVREGRSVML TSHS MEECEALCSRLA IVMN GFRCLGSPQHLKGRF	2016
		*** . * . *****	*****

SEQ2 2039 GAGHTLRLVPPDQPEPAIAFIRITFPGAEELREVHGSRLRFQLPPGRCT 2088
SEQ5 2017 AAGHTLRLVPAARSQPAAAFVAAEFPGSELREAHGGRRLRFQLPPGRCA 2066
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SEQ2 2089 LTRVFRELAAQGRAHGVEDFSVSQTTLEEVFLYFSKDQGEEEESSRQ-EA 2137
SEQ5 2067 LARVFGELAVHGAEHGVEDFSVSQTMLEEVFLYFSKDQGKDEDTEEQKEA 2116
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SEQ2 2138 EEEEVSKPGRQHPKRVSRFLEDPSVETMI 2167
SEQ5 2117 GVGVDPAPGLQHPKRVSQFLDDPSTAETVL 2146
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